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54 TCG	108 CAT H	_	216 GGC G	270 GTG V	324 TTT F	378 AAG K
CTG GCC	AAA K		CAC	CGA	GAC	3TT J
	GAA	ATG	CCT	TCC	GAA	I AT
36 45 CTA GAT CCC TCC	99 ACT T	153 AGC S	207 ATT I	261 GGA G	315 AAC N	369 GAA 1
သသ	AAA K	AGC	GGT G	CT	ACT .	GTG (
GAT	CCC		GTA V	CAT H	ACT	TCA (
36 CTA	90 AAA K	144 GGC G	198 GTT V	52 GT	306 ATC	360 CTT 3
AAT	GAT D	TAT Y	GCT A	3AT D	CAC	GAG (E
CTC	TTG	CGT R	ACA T	ľ.	GAA (	CTT (
27 GCT	81 TTT F	135 TTA L	189 CAC H	243 GTT V	297 TTA (	351 GCT ( A I
TCA	GCT A	GGT G	GCA A	GCA	TTA T	TCT C
909	GGT G	AAT N	GAT D	Tyly	CAT 1	GGA 1 G
18 CAG	72 ATG	126 GGG G	180 GAA E	234 TYT F	288 ACA ( T	342 TCA (
TAT	AAC	GCT A		Ą	<b>ĕ</b>	Æ
ACC	CAA AAC AT	GGT GCT	GAA ATG E M	TGG	279 AT TAC TGC TC Y C. S	GGA A
9 TGT	63 TTG	117 CAT H	171 GTG V	225 GAC ' D	279 FAC :	333 GCT ( A (
TAT	TTA	GCT A	AGA R	GAA (E	AAT 7	GCA G
9 NNA TAT TGT ACC TAT	GAC TTA	AAT	TGG W	TTG ( L I	GCA AAT A N	AGG G R A
		. —	÷	C H	) R	A K

432 TCA S		54.0 TAT Y	594 AGG R	648 AAT N	702 GAT D	756 ATT I
: 1414 F		CTG		GTT	STT	3AA 1
, AAC N		GTT V	AAT N	CGT R	GT	'AT
423 CGT R			385 GC	639 CAA Q	693 AAG K	747 3TY 1
ATG M		CGT R	CCT	ATA [	'AC	AG
TAC	GCA	TCA S	<b>A 1</b>		AT	Ę
414 GAA E	46 AC	522 GA1 D	576 CAC H	630 3TG 7	584 PAT	38 34G
GAT	TC	වි ව	SAT O	SGC ,	AC	CA.
ATT	9 T GGT G	TGI	CAG	25	366 3	CT (
40 AA K	45 AGʻ S	51. AA( N	567 ACC T	621 GGA G	675 CTG	729 3TT 1
TTG	AGG R	ATC I	TCT	GCA A	GCT (	TT
TTT	GAC D	T'I'I F	Tyr		CGT R	CAA (
396 GGA G	450 ATG M	504 TAC Y	558 TGC C	512 2AA 2	566 CT	720 GAA ( E (
ACT	999 9	ATC I	GTC V	CGA ATC ( R I (	GTA V	ACA (T
AGA R	AAC N	CAT H	caa Q	CGA R	GCA (	CCA 7
387 ATC I	441 AGA R	495 AAG K	649 GA	603 GAG E	657 TTA (	711 GGC ( G I
GGT	CTC L	CCT P		AAG K	TCA	AAG ( K (
AAT N	GAC D	TCA S	AGG R	GAA	GGT G	7 299 7 299
		•			- 0	

# FIGURE 1B

810 GAT D	864 TCT S	918 3GA 3	972 3TC 7	126	80 AT	089 1098 1107 1116 1125 1134 CGC ATC TTG TCT GCA GAA AAT ATC CCA AAT TTG CCT CCT GGG GGA GGT R I L S A E N I P N L P P G G G
TGG W	GTA	AAG (	2 AAG (	10 CA C	10 CC C	11 SA G
ATC	GAG	CAC ,	CCC 7	SAA T	TT G	ტ უ
801 GGG G	855 CTT L	909 TTA L	963 GCT (	017 PTG (	371 3AT C	.25 .cr g
GAT D	AGG R	TGT	AAT N	CAC 1	1( CCT ( P I	11 CT C
TGT	TCT	ACT T	TCA	AAG K	ATG (	I'TG (
792 GCT A	846 AAA K	900 GAC D	954 TYT F	.008 GAT D	062 GGA .	116 AAT 3
TTG	GTT V	GTG V	TGC	1 TTG L	1 GAA E	CCA P
ATC I	TAT	GTA V	GTT V	GAG E	GAG	ATC
783 ATC	837 GAA	891 TGG W	945 CTA L	999 TCA S	.053 GGC G	107 AAT N
\ T'T'\ F	TGT C	AAT N	GTA V	GAT D	1 TCT S	1 GAA E
GAP	CTC	TGC	ATT I	AAA K	AAG K	GCA A
774 3 GAT D	828 GAG E	882 GTG V	936 AGT S	990 AAA K	.044 GAG E	098 TCT S
GAG E	GAG	AAT N	ATG M	GTG V	1 ATG M	1 TTG L
GAA	AAT	GAA E	AAC N	GCG A	ATT I	ATC I
765 TTA AGA GCA L R A	819 3 AGT S	873 CTG L	927 GAT D	981 GAA E	1035 GAG E	089 CGC R
AGA R	ATG M	8 T GAC C D L	CGA R	GAT. D	1 GAA E	1089 ATG CGC M R
TTA L	GTT	GAT D	AGT CGA S R	981 TCA GAT GAA S D E	GTT GAA V E	GTC

### FIGURE 10

	•					•
1188 A CAT H	1242 A AAA K	1296 . CAA Q	1350 GAA E	1404 AAC N	1458 TTA L	.512 CTT
, CC2	5 755	CGA R	GAA E	GGA	GAA E	1 rga
AAT N	CAG	TAC Y	GGA	GCT	GCT	ATA '
1179 \ CTG	1233 TCA S	1287 AAC N	1341 GAG E	.395 GAT D	449 CTT L	503 AAA K
' AGA R	GGA	GGA G	GTA	1 AGT S	1 GGT G	1 3AA , E 1
AGT	AGT S	AGG R	AAA K	AAC N	AGT S	GGT (
1170 r tat Y	1224 3 GAA E	1278 CAT H	1332 GCT A	1386 TCG S	440 GAA E	494 AGT ( S (
, GT	GAC	AAT N	CTA L	TCT	TCA S	1 ATG M
GCT	GCA	ATT I	AGG R	ACT	GAA E	AAG K
1161 ' GAA E	1215 GAA E	1269 AGA R	1323 TAC Y	.377 GCT A	431 ACT T	485 ACA 7
ATT	GAT D	ATG	AGT	1 ACA T	1 CAT H	366 1
GTT	TCC	CAA Q	ACT	GCC A	AGC	SCA (
1152 AAT N	1206 GCC A	1260 AGG R	.314 CTG L	368 GCT A	422 GAA E	476 3AT (
CGT R	999 9	CTC	ATG M	CCA P	CAG	1 3AA (
AAG K	GAT D	GCT A	GAG	GAA E	ATG	AAT (
1143 GGC G	1197 AGT S	.251 GAA E	305 GAG E	359 GCT A	413 ACA F	167 PCT 1
GCT	GAA	GTG V	1 CTG L	CCT P	1. 3TG /	14 AGC 1
CTT GCT GGC AAG CGT AAT GTT ATT GAA GCT GTT TAT AGT AGA CTG AAT CCA CAT L A G K R N V I E A V Y S R L N P H	AGA R	TTG	CTT	AGC (	CCA (	GAC A D S

#### FIGURE 11

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TCC TIT TIG GTA ATA TIT TIG TGA TCT TIG ATG GIT TIT AAC CTA GGA AGT GTA TAT TTG AAT CTT GGA AAA CTA GTT TTA TTT TGT TAT GCA TTT ATA TAA CTG 

ATG

TTA TAT TCA GAT AGC CTT GTT TTT TAA AAA GGC CTT TGC ATA CAC CTT TAT GAG 

TGA AAT TAT ATG TCA ATA GTG TAA AAT TGA CTA TTT ATA GTA CTA TGG ATT TAA 

TTT CAC ATT GTA TGC CAG AAA TTA GGC TAC CAA TTA TGA ATT AAA GTC AGT AGT 

TAA ATT AAT ACT AGA TAG AAT TAG AAA TTT TGA TTA GAG AGA TTA TGC TAT ATT 

TGC TTC ATA TTA TTT TAC CTA TTA GAA TTA TAC GTT AAT GTA GAA AAA CTT 

### FIGURE 1E

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Title: HUMAN PROTEIN PHOSPHATASE Serial No.: To Be Assigned

1935 1944 TYT AAT AAT TYT AGT	1989 TCA AGG ATG ATA ATT	2034 2043 2052 ATC TCT GTA GTT TTA TTT GAA	2106 CTT TAT GAA
1935 TTT AAT	1989 TCA AGG	2043 GTT TTA	2097 CTC TGT ATT (
TTC	CTT	GTA	CTC
1917 TAA TAA ATT TAT GTT TYC	1980 ATT TYT CTT	2034 ATC TCT	2088 TCC TTT
ATT	GTA	TAC	I'I'I
1917 TAA TAA	1971 TGG CCT	2025 ATA TYT TAC	2079 GGC TAT TYT
TTG	TGC	IAIA	TGT
TAG TTA GCT TTG	1962 AAT GGC TGA TGC	2016 GAT TYG	2070 TGG ATG
		Tyry	TAT
GTA CAC TCA	1953 FCT TCA AAG	2007 TGT GTG TTG	2061 GTT GTG AGA
GTA	TCT	TGT	GTT

TGT ATA GCC TAA CTA CAC ACA TCA AAA TGT ATG TCA ACC AAG TGT TTA GAA TGA 

ACA TAA CTT TTG AAA AAC CTA TGT ATT ATT CAT ACA GCT TTG GTT TGT ATA TTC

AAT TAT AAG TGT TTA AGT CCA AAT AAA GCA TGT GAT GTG GAA TAA TCA AAA AAA 

## FIGURE 1F

FIGURE 1G

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Title: HUMAN PROTEIN PHOSPHATASE Serial No.: To Be Assigned GI 452526 452526 GI 247927 GI 247927 452526 452526 247927 247927 247927 452526 13177 GI GI G 댕 GI GI Ŀ Ŀ F 0 0 G R R 2 K Ø Z Z Z ZZ Z S S S Q Q 王 H  $\approx$  $\alpha$ 2  $\simeq$  $\mathbb{R}$ 2 > H 田 团 口 团 C Σ Σ C Ø K Ø  $\mathbf{\Sigma}$ Ц V L H 口 J Ц 田 H S  $\Omega$ S > H 王 H K G G G K H K K 24 Z Z Z 3 S S S × S S > S > G G G  $\circ$  $\mathcal{C}$ П Ω Q  $\mathbb{R}$  $\approx$  $\alpha$ Ø 0  $\succ$ Ŀ Ŀ G G G Ŏ Ŏ 0  $\mathbf{\Sigma}$ Σ  $\mathbf{\Sigma}$ Z Z Z G G G  $\circ$ C O S ß Ø Ø K Z Z Z Σ Σ Σ Ω Ü S > > > >  $\alpha$  $\mathbb{A}$ K > > J 24 24 Н Ţ, [T. Ŀ S S S G G 9 S S S G G S  $\succ$ 5 5 G G G Ö Z T Ţ M G G Ö NVK K  $\alpha$ 24 K K K X H 工 Z A A 团 П J  $\Xi$ > H > Ţ 뇌 Z Z Z G G S G G S Z S Д Д  $\alpha \alpha o$ S Z Z Z Ω Q Q **1**1 Ы  $\Omega$ 田 S  $\vdash$ Н G G G  $\succ$ > > > >  $\simeq$ K L S S K K K > > > S  $\mathbf{z}$ Σ H H 回 Ŏ G G G Ø Ø K Р Д > > >  $\mathbf{\times}$ [I]  $\Xi$ 工 H Ţ, H 国<sub>,</sub> 99 G घ घ 团 T T K Ø Ė J Œ, ĹŢ, ļ > > Σ > Z Z Z  $\Omega$ ¥ S K K N K Д A Д Б G 工  $\Xi$ 3 3 3 SI S Ŧ Z Ę Z × ပ်ပြ × Z O SSS  $\mathcal{O}$ G [I] 团 团 S S S0 G Д Ω DK ᆸ J, ×  $\Omega$ S × × × G G G ଠା Ω K  $\alpha$ 工  $\Xi$ X Ξ  $\circ$ Q, H ব H H ОО Q  $\succeq$ Д Д K Ø ΣΣ Ø  $\alpha$ Σ 0  $\vdash$  $\alpha$ G Ŋ G Ę E Ω 二  $\Box$ G Ŋ G Œ, Z Z Z SSS Ŀ > >Д >  $\aleph$ 24 נבי נבי K > > > H ध ध Ц П  $\circ$ C G G G C Ø K K Z Ω Z Z Q > > > ļ Σ ₽ ₽ E S Ø  $\Omega$ S Ŏ 0 K 161 121 161 121 4 网络网络

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		Serial No.:	To Be Assigned		
13177 GI 247927 GI 452526	13177 GI 247927 GI 452526	13177 GI 247927 GI 452526	13177 GI 247927 GI 452526	13177 GI 247927 GI 452526	13177 GI 247927
241 A C D G I W D V M S N E E L C E Y V K S R L E V S D D L E N V C N W V V D T C L 241 A C D G I W D V M S N E E L C E F V M S R L E V S D D L E N V C N W V V D T C L 241 A C D G I W D V M S N E E L C E F V K S R L E V S D D L E N V C N W V V D T C L	281 HKGSRDNMSIVLVCFSNAPKVSDEAVKKDSELDKHLESRV 281 HKGSRDNMSIVLVCFANAPKVSDEAVKRDLELDKHLESRV 281 HKGSRDNMSVVLVCFSNAPKVSEEAVKRDSELDKHLESRV	321 EEIMEKSGEEGMPDLAHVMRILSAENIPNLPPGGLAGKR 321 EEIMOKSGEEGMPDLAHVMRILSAENIPNLPPGGGLAGKR 321 EEIMOKSGEEGMPDLAHVMRILSAENIPNLPPGGGLAGKR	361 NVIEAVYSRLNPHRESDGASDEAEESGSOGKLVEALROMR 361 NVIEAVYSRLNPNKDNDGGAGD	401 INHRGNYROLLEEMLTSYRLAKVEGEESPAEPAATATSSN 383	441 SDAGNPVTMOESHTESESGLAELDSSNEDAGTKMSGEKI 390

### FIGURE 21

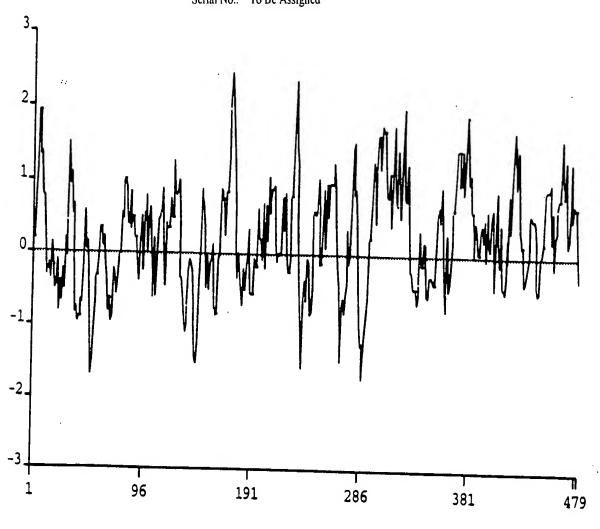


FIGURE 3A

